

**AMENDMENTS TO THE CLAIMS**

**LISTING OF CLAIMS:**

*This listing of claims will replace all prior versions and listings of claims in the application:*

1. (Original) An isolated peptide comprising the amino acid sequence QA(Q/E)GQLV or functional equivalents thereof, wherein said peptide selectively homes to TNF receptor(s) of the vasculature of a heart.

2.-17. Cancel

18. (Original) An isolated peptide comprising the amino acid sequence ARRGQAV or functionally equivalent thereof, wherein said peptide preferentially homes to BDNF receptor(s) of the vasculature of a heart.

19.-32. Cancel

33. (Original) An isolated peptide comprising the amino acid sequence G(R/W)RFIRV or functional equivalent thereof, wherein said peptide preferentially homes to BDNF receptor(s) of the vasculature of a heart.

34.-51. Cancel

52. (Original) A conjugate comprising a peptide according to claim 1 and a functional moiety, wherein said peptide selectively homes to TNF receptor(s) in the vasculature of a heart.

53.-65. Cancel

66. (Original) A conjugate comprising a peptide according to claim 18 and a functional moiety, wherein said peptide preferentially homes to BDNF receptor(s) of the vasculature of a heart.

67.-74. Cancel

75. (Original) A conjugate comprising a peptide according to claim 33 and a functional moiety, wherein said peptide preferentially homes to BDNF receptor(s) of the vasculature of a heart.

76.-85. Cancel

86. (Original) A method for determining a young heart or young areas of a heart vasculature in a mammal comprising:  
a) administering a peptide comprising the amino acid sequence QA(Q/E)GQLV or functionally equivalent modifications thereof, conjugated to a detectable marker, wherein said first peptide selectively homes to TNF receptor(s) in a vasculature of the heart; and  
b) detecting the marker;  
wherein a disproportionately high binding of QA(Q/E)GQLV is a young heart or young areas of a heart vasculature.

87.-91. Cancel

92 (Original) A method for determining an old heart or old areas of a heart vasculature in a mammal comprising:  
a) administering a peptide comprising the amino acid sequence ARRGQAV or G(R/W)RFIRV or functionally equivalent modifications thereof, conjugated to a detectable marker, wherein said first peptide selectively homes to BDNF receptor(s) in a vasculature of the heart; and  
b) detecting the marker;

wherein a disproportionately high binding of ARRGQAV or G(R/W)RFIRV is an old heart or old areas of a heart vasculature.

93.-101. Cancel

102. (Original) A method for determining the condition of a vasculature of a heart in a mammal comprising:

- a) administering a first peptide comprising the amino acid sequence QA(Q/E)GQLV or functionally equivalent modifications thereof, conjugated to a first detectable marker, wherein said first peptide selectively homes to TNF receptor(s) in the vasculature of the heart;
  - b) administering a second peptide comprising the amino acid sequence ARRGQAV or G(R/W)RFIRV or functionally equivalent modifications thereof, conjugated to a second detectable marker, wherein said second peptide homes to BDNF receptor(s) in the vasculature of the heart; and
  - c) detecting the first and second marker;
- wherein a disproportionately high ratio of binding of the first peptide to the second peptide indicates a young heart or young areas of the heart vasculature or wherein a disproportionately low ratio of binding of the first peptide to the second peptide indicates an old heart or old areas of the heart vasculature.

103.-115. Cancel

116. (Currently Amended) A method for delivering a functional moiety to a young heart vasculature in a mammal, the method comprising administering a conjugate of claim 48 52.

117.-126. Cancel

127. (Currently Amended) A method for delivering a functional moiety to a old heart vasculature in a mammal, the method comprising administering a conjugate of claim 66 or ~~claim 75~~.

128.-138 Cancel

139. (Original) A method for discovering mimics of amino acid sequence QA(Q/E)GQLV or functionally equivalent modifications thereof, comprising:

- a) determining a three-dimensional structure of said sequence;
- b) identifying compounds comprising said structure; and
- c) determining the capacity of said compounds for selective homing to TNF receptor(s) in a heart vasculature of a mammal;

wherein compounds which selectively home to TNF receptor(s) in the vasculature of the heart are mimics.

140.-146. Cancel

147. (Original) A method for discovering mimics of amino acid sequence ARRGQAV or G(R/W)RFIRV or functionally equivalent modifications thereof, comprising:

- a) determining a three-dimensional structure of said sequence;
- b) identifying compounds comprising said structure; and
- c) determining the capacity of said compounds for homing to BDNF receptor(s) in a heart vasculature of a mammal;

wherein compounds which home to BDNF receptor(s) in the vasculature of the heart are a mimics.

148.-156 Cancel

157. (New) A method for delivering a functional moiety to a old heart vasculature in a mammal, the method comprising administering a conjugate of claim 75.